

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 10/579,290
Source: IFWP
Date Processed by STIC: 2/1/07

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

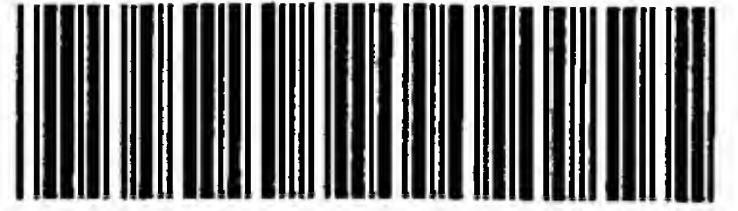
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Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

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1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/579,290

DATE: 02/01/2007

TIME: 11:51:19

Input Set : A:\pto.ss.txt

Output Set: N:\CRF4\02012007\J579290.raw

4 <110> APPLICANT: OncoMab GmbH
 6 <120> TITLE OF INVENTION: Human monoclonal antibody
 C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/579,290
 C--> 8 <141> CURRENT FILING DATE: 2006-05-15
 W--> 0 <130> FILE REFERENCE:
 8 <150> PRIOR APPLICATION NUMBER: EP - 03 026 161.4
 9 <151> PRIOR FILING DATE: 2003-11-14
 11 <160> NUMBER OF SEQ ID NOS: 4

ERRORED SEQUENCES

13 <210> SEQ ID NO: 1
 14 <211> LENGTH: 96
 15 <212> TYPE: DNA
 16 <213> ORGANISM: Homo sapiens
 18 <220> FEATURE:
 19 <223> OTHER INFORMATION: Amino acid sequence of the variable region of the light chain

(VL) of
 20 antibody
 21 SAM-6

23 <400> SEQUENCE: 1
 25 Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly
 26 1 5 10 15
 29 Gln Thr Ala Ser Ile Thr Cys Ser Gly Asp Lys Leu Gly Asp Lys
 30 20 25 30
 33 Tyr Ala Cys Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu
 34 35 40 45
 37 Val Ile Tyr Gln Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg
 38 50 55 60
 41 Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser
 42 65 70 75
 45 Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp
 46 80 85 90

For AMINO
 Acid
 Sequences.

49 Asp Ser Ser Ile Val Val
 50 95

113 <210> SEQ ID NO: 2
 114 <211> LENGTH: 110
 115 <212> TYPE: DNA
 116 <213> ORGANISM: Homo sapiens
 118 <220> FEATURE:
 119 <223> OTHER INFORMATION: Amino acid sequence of the variable region of the heavy chain (VH) of antibody

121

SAM-6

See error explanation on page 1,

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Input Set : A:\pto.ss.txt

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123 <400> SEQUENCE: 3

125 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly
 126 1 5 10 15
 129 Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
 130 20 25 30
 133 Ser Tyr Ala Met His Trp Val Arg Glu Ala Pro Gly Lys Gly Leu
 134 35 40 45
 137 Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr
 138 50 55 60
 141 Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser
 142 65 70 75
 145 Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
 146 80 85 90
 149 Thr Ala Val Tyr Tyr Cys Ala Arg Asp Arg Leu Ala Val Ala Gly
 150 95 100 105

153 Lys Thr Phe Asp Tyr

154 110

167 <210> SEQ ID NO: 4

168 <211> LENGTH: 330

169 <212> TYPE: DNA

170 <213> ORGANISM: Homo sapiens

172 <220> FEATURE:

173 <223> OTHER INFORMATION: Nucleotide sequence of the variable region of the heavy
 chain (VH) of

174 antibody

175 SAM-6

177 <400> SEQUENCE: 4

179 cag gtg cag ctg gtg gag tct ggg gga ggc gtg gtc cag cct ggg 45
 180 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly
 181 1 5 10 15
 184 agg tcc ctg aga ctg tcc tct gca gcc tct gga ttc acc ttc agt 90
 185 Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
 186 20 25 30
 189 agc tat gct atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg 135
 190 Ser Tyr Ala Met His Trp Val Arg Glu Ala Pro Gly Lys Gly Leu
 191 35 40 45
 194 gag tgg gtg gca gtt ata tca tat gat gga agc aat aaa tac tac 180
 195 Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr
 196 50 55 60
 199 gca gac tcc gtg aag ggc cga ttc acc atc tcc aga gac aat tcc 225
 200 Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser
 201 65 70 75
 204 aag aac acg ctg tat ctg caa atg aac agc ctg aga gct gag gac 270
 205 Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
 206 80 85 90
 209 acg gct gtg tat tac tgt gcg aga gat cgg tta gca gtg gct ggt 315
 210 Thr Ala Val Tyr Tyr Cys Ala Arg Asp Arg Leu Ala Val Ala Gly
 211 95 100 105

E--> 214 aaa act ttt gac tac

215 Lys Thr Phe Asp Tyr

PLS use
 type PRT
 for amino
 acid
 sequences,

330 ← Insert

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/579,290

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Input Set : A:\pto.ss.txt

Output Set: N:\CRF4\02012007\J579290.raw

W--> 216

110

VERIFICATION SUMMARY

DATE: 02/01/2007

PATENT APPLICATION: US/10/579,290

TIME: 11:51:20

Input Set : A:\pto.ss.txt

Output Set: N:\CRF4\02012007\J579290.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
L:50 M:301 E: (44) No Sequence Data was Shown, SEQ ID:1
L:50 M:252 E: No. of Seq. differs, <211> LENGTH:Input:96 Found:0 SEQ:1
L:154 M:301 E: (44) No Sequence Data was Shown, SEQ ID:3
L:154 M:252 E: No. of Seq. differs, <211> LENGTH:Input:110 Found:0 SEQ:3
L:214 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:330 SEQ:4
L:216 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4